SEQUENCE LISTING

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<110> Anderson, Christen M.
      Clevenger, William
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Lys Arg Glu Lys Ala Glu Glu Asp Arg Tyr Phe Arg Glu Lys Thr Arg 50 55 60	
Glu Gln Leu Ala Ala Leu Lys Lys His His Glu Asp Glu Ile Asp His	
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Glu Gln Leu Ala Ala Leu Arg Lys His His Glu Asp Glu Ile Asp His
                    70
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     Thr Arg Glu Gln Leu Ala Ala Leu Lys
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     <210> 26
     <211> 17
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     <213> Artificial Sequence
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     <223> Polypeptide consisting of amino acids 42-58 of the
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Leu Ala Ala Leu Lys Lys His His Glu Asp Glu Ile Asp His His Ser
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    Lys
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                 20
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      sequence.
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Glu Asp Arg Tyr Phe Arg Glu Lys Thr Arg Glu Gln Leu Ala Ala Leu
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Lys Lys
<210> 30
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Glu Asp Arg Tyr
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            20
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           sequence.
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                                          10
     Tyr Phe Arg Glu
William Marie
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                                          10
                                                               15
     Phe Arg Glu Lys
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     <210> 35
     <211> 20
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     <213> Artificial Sequence
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    <400> 35
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LICESSIS OBEVIL
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1
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Arg Glu Lys Thr
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<210> 36
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Lys Thr Arg Glu
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Thr Arg Glu Gln
            20
```

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<220>
          Ξ
          Appendix of the control of the contr
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<223> Synthetic peptide fragment derived from rat IF1
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Leu Ala Ala Leu
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Ala Ala Leu Lys
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Ala Leu Lys Lys
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                 5
                                     10
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     Ser Ile Arg Glu Ala Gly Gly Ala Phe Gly Lys Arg
      1
<210> 47
     <211> 13
     <212> PRT
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     <213> Artificial Sequence
į.
     <220>
     <223> Synthetic peptide fragment derived from rat IF1
U
           sequence.
2
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Ser Ile Arg Glu Ala Gly Gly Ala Phe Gly Lys Arg Glu
     1
                      5
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     <210> 49
     <211> 15
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     <223> Synthetic peptide fragment derived from rat IF1
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<210> 46 <211> 12 <212> PRT

<213> Artificial Sequence

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      1
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     <211> 17
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L.
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     Glu
     <210> 52
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     Glu Asp Arg Tyr Phe
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1
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Glu Asp Arg Tyr Phe Arg
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                                     10
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            20
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Glu Asp Arg Tyr Phe Arg Glu Lys
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                                     10
Glu Asp Arg Tyr Phe Arg Glu Lys Thr
            20
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Glu Asp Arg Tyr Phe Arg Glu Lys Thr Arg Glu
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Ser Ile Arg Glu Ala Gly Gly Ala Phe Gly Lys Arg Glu Lys Ala Glu

Glu Asp Arg Tyr Phe Arg Glu Lys Thr Arg Glu Gln Leu

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20
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Glu Asp Arg Tyr Phe Arg Glu Lys Thr Arg Glu Gln Leu Ala Ala Leu

sequence.

<400> 63

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Lys
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Gly Gly Gln Gln Met Gly Arg Asp Leu Tyr Asp Asp Asp Asp Lys Asp
                                 25
                                                     30
Pro Ser Ser
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Met Arg Val Leu Gln Thr Arg Gly Phe
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Pro Ser Ser Gly Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Gly
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Met Ala Gly Ser Ala Leu Ala Val Arg Ala Arg Leu Gly Val Trp Gly
Met Arg Val Leu Gln Thr Arg Gly Phe Ser Ile Arg Glu Ala Gly Gly
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                                      75
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Lys Thr Arg Glu Gln Leu Ala Ala Leu Lys Lys
           100
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ggtgtctggg gtatgagggt cctgcaaacc cgaggcttct ccatccgaga agctggtggg 240
gccttcggga aacgagagaa ggctgaagag gatcggtact tccgagagaa gactagagag 300
cagctggctg ccttgaagaa g
                                                               321
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